```
gtccaaggtc cagaaagcaa aaagggtgcc ccgtttctcc tgggaaggct tccccgtcag       2460

tgatttctgt aaccggaccc tgccctgaca cagcgtcatt ggactaccca gcagacagta       2520

gactccactc taaacccgct tcttgcggtc agttgctgtc cttcagtgtg tgtaagcagt       2580

ggccagacag cacccttggg tgtcatttca agactctctc accttggtct gctttacgtt       2640

tggtttgatt tggtttgttc tggtttttga dacgaggcct ttcactggaa cctggcactc       2700

agtatttaga ctgcccagcc agctagcctc agagaatgca tctgcgtatg cttgcctggc       2760

gctggaattc ggtgcacatg gctttgatgt gtaccgggga tcagacacag atgtttcatg       2820

agtgcagtgc atgcctgtta gtggtagagc tc                                    2852
```

<210> 8
<211> 41
<212> PRT
<213> Homo sapiens

<400> 8

```
Met Leu Arg Ala Leu Asn Arg Leu Ala Ala Arg Pro Gly Gly Gln Pro
1               5                   10                  15


Pro Thr Leu Leu Leu Leu Pro Val Arg Gly Pro Arg Pro Arg Ser Phe
            20                  25                  30


Ser Ala Pro Phe Ser Ser Gln Asp Ser
            35                  40
```

<210> 9
<211> 20
<212> DNA
<213> Artificial

<220>
<223> SYNTHETIC PRIMER

<220>
<221> misc_feature
<222> (1)..(20)
<223> SYNTHETIC PRIMER

<400> 9
ttcacaccac tctgtcgaac   20

<210> 10
<211> 20
```

```
<220>
<221>  misc_feature
<222>  (1)..(19)
<223>  construct

<400>  36
aggcataaaa aagtcggtc                                                    19


<210>  37
<211>  15
<212>  PRT
<213>  Unknown

<220>
<223>  conserved sequence


<220>
<221>  DOMAIN
<222>  (1)..(15)
<223>  conserved sequence

<400>  37

Pro Arg Pro Arg Ser Phe Ser Ala Pro Phe Ser Ser Gln Asp Ser
1               5                   10                  15
```